

1 **Title**

2 Mass coral bleaching causes biotic homogenization of reef fish assemblages

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4 **Running head**

5 Coral bleaching causes biotic homogenisation

6

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18

19 **Key words**

20 *Climate change; coral reefs; assemblage structure; coral species composition; functional*

21 *redundancy; traits; scales; beta diversity*

22

23 **Paper type**

24 Primary research article

25 **Abstract**

26

27 Global climate change is altering the composition of ecosystems due to non-random species  
28 turnover, typically characterized by the loss of specialist species and increasing similarity of  
29 biological communities across spatial scales. As anthropogenic disturbances continue to alter  
30 species composition globally, there is a growing need to identify how species responses  
31 influence the establishment of distinct assemblages such that management actions may be  
32 appropriately assigned. Here, we use trait-based analyses to compare temporal changes in  
33 five complementary indices of reef fish assemblage structure among six taxonomically  
34 distinct coral reef habitats exposed to a system-wide thermal stress event. Our results  
35 revealed increased taxonomic and functional similarity of previously distinct reef fish  
36 assemblages following mass coral bleaching, with changes characterized by subtle but  
37 significant shifts in dominant fish taxa towards small-bodied, algal-farming habitat  
38 generalists. Furthermore, whilst the taxonomic or functional richness of fish assemblages did  
39 not change across all habitats, an increase in functional originality indicated an overall loss of  
40 functional redundancy. We also found that pre-bleaching coral composition better predicted  
41 changes in fish assemblage structure than the magnitude of coral loss. These results  
42 emphasise how measures of alpha diversity can mask important changes in the structure and  
43 functioning of ecosystems as assemblages reorganize. Our findings also highlight the role of  
44 coral species composition in structuring communities and influencing the diversity of  
45 responses of reef fishes to disturbance. As new coral species configurations emerge, their  
46 desirability will hinge upon the organisation of associated species and their capacity to  
47 maintain key ecological processes in spite of ongoing disturbances.

48

49 **Introduction**

50

51 Pervasive anthropogenic disturbances have altered the structure of ecological communities  
52 and the functioning of ecosystems, primarily through habitat change, exploitation, and  
53 pollution (Cardinale et al., 2012; Ellis et al., 2013). Species extirpations and introductions  
54 modify the composition of local assemblages, often reducing biodiversity within (alpha  
55 diversity;  $\alpha$ ), and among (beta diversity;  $\beta$ ) communities (Dornelas et al., 2014; McGill,  
56 Dornelas, Gotelli, & Magurran, 2015). This resulting species turnover typically occurs non-  
57 randomly, often varying with organism characteristics such as body-size, physiology, habitat  
58 specialisation and trophic level (McKinney, 1997; Graham et al., 2011). Long-term  
59 observations of trait-filtering across groups such as plants, mammals, birds, and fishes,  
60 indicate declines in specialist species in response to past and current global changes (Clavel,  
61 Julliard, & Devictor, 2011). These declines have caused increasing taxonomic and functional  
62 similarity (i.e., biotic homogenization) at the community level, threatening ecosystem  
63 functioning and resilience (Elmqvist et al., 2003; Olden, LeRoy Poff, Douglas, Douglas, &  
64 Fausch, 2004; Clavel et al., 2011). Two facets of species biodiversity are expected to confer  
65 ecosystem resilience to disturbance during ecological reorganization: (i) functional  
66 redundancy (the capacity of one or more species to functionally compensate for the loss of  
67 another), and (ii) response diversity (the diversity of species responses to environmental  
68 change within functional groups such that functions may be maintained) (Walker, 1992;  
69 Elmqvist et al., 2003; Nyström, 2006; Nash, Graham, Jennings, Wilson, & Bellwood, 2016 ).

70

71 We focus on coral reefs, one of the world's most biodiverse but threatened ecosystems  
72 (Hughes et al., 2017), to assess how species response diversity to disturbance influences  
73 ecological organisation across scales (within and across distinct reef habitats). Climate-  
74 induced coral bleaching represents the foremost threat to coral reefs, with severe thermal

75 stress events causing widespread coral loss, altering the structure of coral reef habitats and  
76 the composition of reef-associated species (Graham et al. 2006; Hoegh-Guldberg et al. 2007;  
77 Pratchett, Hoey, Wilson, Messmer, & Graham, 2011). Mass bleaching has been shown to  
78 cause local extirpations of coral reef fishes with corresponding declines in fish species  
79 richness and total abundance (Wilson, Graham, Pratchett, Jones, & Polunin, 2006). However,  
80 reef fish and coral species responses to extreme thermal stress can be highly variable (Wilson  
81 et al., 2006; McClanahan, 2017), likely due to inherent differences in the response of corals  
82 to ocean warming (Pandolfi, Connolly, Marshall, & Cohen, 2011), as well as differential  
83 trait-vulnerability of fish species associated with distinct coral habitats (Graham et al., 2011;  
84 Richardson, Graham, Pratchett, & Hoey, 2017). Sustained ocean-warming increases the risk  
85 of further mass bleaching events (Hughes et al., 2017). However, how distinct species  
86 assemblages will vary in their susceptibility to change is not well understood.

87

88 There is a growing need to understand how different configurations of species on coral reefs  
89 respond to disturbance and contribute to ecosystem function (Graham, Cinner, Norström, &  
90 Nyström, 2014), such that coral reef management efforts may be allocated appropriately.  
91 Previously unseen disturbance-mediated configurations of species are increasingly reported  
92 on coral reefs (Pandolfi et al., 2011; Pratchett, Traçon, Berumen, & Chong-Seng, 2011;  
93 Bento, Hoey, Bauman, Feary, & Burt, 2016). Increasing selective disturbances are causing  
94 shifts in coral species dominance, as well as overall declines in species diversity (Aronson,  
95 Macintyre, Wapnick, & O'Neill, 2004; Pratchett et al., 2011; Alvarez-Filip, Carricart-  
96 Ganivet, Horta-Puga, & Iglesias-Prieto, 2013). These altered configurations are predicted to  
97 persist into the future due to taxa specific responses to perturbations, recovery potential, and  
98 capacity for rapid adaptation under projected climatic changes of global warming and ocean  
99 acidification (Pandolfi et al., 2011). The biological and physical structure of coral reef

100 habitats exerts considerable influence over the structure of reef fish assemblages (Luckhurst  
101 & Luckhurst, 1978; Roberts & Ormond, 1987; Friedlander & Parrish, 1998; Darling et al.,  
102 2017), important ecosystem processes such as herbivory (Cvitanovic & Hoey, 2010),  
103 predator-prey dynamics (Hempson et al. 2017), and other intraspecific and interspecific fish  
104 species interactions (Kok, Graham, & Hoogenboom, 2016). Indeed, shifts in coral  
105 assemblage composition have also shaped novel reef fish assemblages, where despite  
106 regaining pre-disturbance coral cover in some instances, populations of some fish species  
107 show little evidence of recovery generations later (Bellwood et al., 2012; Berumen &  
108 Pratchett, 2006).

109

110 Here we assess the susceptibility of distinct species configurations to the impacts of mass  
111 coral bleaching to elucidate the influence of pre-disturbance coral species composition on the  
112 resilience of tropical reefs, and consider the effects of response diversity on ecological  
113 organisation. Specifically, we use trait-based approaches to compare the taxonomic and  
114 functional structure of reef fish assemblages among six taxonomically distinct coral reef  
115 habitats seven months before, during, and six months after severe coral bleaching at Lizard  
116 Island, in the northern Great Barrier Reef, Australia.

117

## 118 **Materials and methods**

119

### 120 *Study sites*

121

122 Lizard Island (14°41'S, 145°27'E) is a granitic mid-shelf island located 30 km offshore from  
123 mainland Australia, in the northern section of the Great Barrier Reef. The island is largely  
124 encircled by well-developed fringing reefs and an extensive lagoonal system of patch reefs.

125 In 2016, the northern Great Barrier Reef (including Lizard Island) was subject to extreme  
126 temperature stress, with peak daily sea surface temperatures reaching 32.8 °C in February  
127 2016, and a high thermal load of >8 degree heating weeks (°C-weeks) by April 2016,  
128 triggering mass coral bleaching (Hughes et al., 2017).

129

130 Benthic and fish assemblages were surveyed at 16 sites on the leeward side of Lizard Island  
131 seven months before, during and approximately six months after the bleaching event, in  
132 September 2015, April 2016, and October 2016 respectively. Sites were randomly selected in  
133 September 2015 within the following constraints: there was >250 m x 5 m of contiguous reef,  
134 adjacent sites were separated by >500 m, and were protected from prevailing south-east swell  
135 with comparable water clarity and geomorphology.

136

#### 137 *Assessment of benthic and fish assemblages*

138

139 Benthic and fish assemblages were surveyed along six replicate 30 m transects at each site,  
140 positioned on shallow (<6 m) reef edges and parallel to the reef–sand interface in each time-  
141 period. Transect position was initially random in September 2015, and the starting position of  
142 individual transects were subsequently identified visually within the same approximate area  
143 (within 1–2 m) by the same surveyors (LR and JE) for the following survey periods.

144

145 The point-intercept method was used to quantify benthic composition and coral bleaching  
146 extent, recording the substrate immediately below the transect tape at 25 cm intervals (120  
147 points per transect). Substratum categories were hard (scleractinian) and soft (alcyonacean)  
148 corals recorded to genus (or species where possible) and growth form, ‘other sessile  
149 invertebrates’ (primarily sponges, giant clams, and ascidians), macroalgae, erect crustose

150 coralline algae, 'dead substrate' (dead coral and pavement), rubble, and sand. Coral bleaching  
151 was estimated for live coral tissue that lay beneath the survey points (recorded in April and  
152 October 2016). A 6-point colour saturation scale based on the CoralWatch colour reference  
153 card was used *in situ* to minimise subjective assessment of bleaching state (Siebeck, Marshall,  
154 Klüter, & Hoegh-Guldberg, 2006). Coral tissue with saturation scores of 1–2 was considered  
155 bleached and 3–6 unbleached to provide an estimate of the percent cover of bleached coral  
156 along each transect. No explicit account was made of bleaching variation within coral colonies.

157

158 The species identity, abundance, and body-size (total length; TL, to the nearest cm) of all non-  
159 cryptic fishes were recorded on transects. Larger, more mobile species (>10 cm TL) were  
160 counted as transects were laid in a 5 m wide belt, and smaller fishes ( $\leq 10$  cm) were surveyed  
161 in a 1 m belt on the return pass. All fish surveys were conducted by a single observer (JE) and  
162 the fish abundances standardised per 150 m<sup>2</sup>. Fish species were classified into functional  
163 entities according to six ecological and morphological species traits: diet, mean observed body-  
164 size (TL; 10 cm size categories), social grouping, mobility within/between reefs, time when  
165 active, and position in the water column (following Mouillot, Graham, Villéger, Mason, &  
166 Bellwood, 2013) (Table S1). Body-size was estimated with the mean observed sizes of each  
167 species in the September 2015 and October 2016 surveys, unless species were observed only  
168 in April 2016 whereby mean size was estimated for that period only.

169

170 *Data analysis*

171

172 Benthic composition

173

174 Pre-bleaching variation in benthic composition among sites was visualized with non-metric  
175 Multidimensional Scaling (nMDS) based on a Bray-Curtis similarity matrix of square root  
176 transformed transect level data. Groupings identified with nMDS were tested using a two-  
177 way nested PERMANOVA (maximum permutations = 9999), using habitat grouping (fixed)  
178 and site (random) as factors. One-way unrestricted permutations of raw data was used for  
179 pairwise comparisons between habitats to allow testing of sufficient permutations, followed  
180 by percentage similarity analysis (SIMPER) to identify substratum categories consistently  
181 contributing to mean similarity within or dissimilarity between habitats (with  
182 similarity/dissimilarity test ratio of >4.0 or 2.0, respectively (Clarke & Warwick, 2001);  
183 Table S2). The percentage of total coral cover (hard and soft coral) that bleached was  
184 compared among habitats using linear mixed-effects analyses (with lme in *nlme*; R  
185 Development Core Team 2016), using habitat as a fixed effect and site as a random effect.  
186 Changes in total percentage coral cover (hard and soft coral) among habitats following the  
187 bleaching event were assessed with habitat and period (and their interaction) as fixed effects  
188 and site (random effect). *Post hoc* multiple comparison Tukey tests were used to identify  
189 where differences occurred.

190

191 Fish assemblage structure

192

193 Trait-based functional diversity was calculated by constructing a principal coordinates  
194 analysis (PCoA) of species pairs based on trait combinations and a Gower distance matrix,  
195 with a square root correction for negative eigenvectors (Legendre & Legendre, 1998). Four  
196 synthetic PCoA axes summarizing species distribution within functional trait space were then  
197 combined with species abundances to calculate three complementary indices of functional  
198 diversity for each transect in all periods: functional richness, functional dispersion, and



199 functional originality (Mouillot et al., 2013; Maire, Grenouillet, Brosse, & Villéger, 2015).  
200 Functional richness is calculated as the proportional convex hull volume occupied by a  
201 species assemblage in multidimensional functional space (PCoA), and represents the range of  
202 unique trait combinations. Functional dispersion measures the abundance-weighted deviation  
203 of species traits in an assemblage from the centroid of all species in functional space, and  
204 reflects the assemblage spread, or variability, of species traits (Laliberté & Legendre, 2010).  
205 Functional originality is calculated as the mean pairwise distance between species in an  
206 assemblage where greater mean distance infers greater functional originality and isolation of  
207 species in functional space. Functional richness and functional dispersion provide presence-  
208 absence and abundance-weighted facets of functional diversity respectively, whilst functional  
209 originality indicates how changes in species abundances modify the functional redundancy  
210 among species (Mouillot et al., 2013). Variation in the functional structure of fish  
211 assemblages through time was assessed using community-weighted transect means (CWM)  
212 of assemblages in each habitat before, during and after bleaching, measured using the four  
213 PCoA axes combined with species abundance values.

214

215 Linear mixed-effects models were used to assess variation in the structure of fish  
216 assemblages in two distinct analyses, both (i) within each habitat across time-periods, and (ii)  
217 among habitats within each time-period. First, within-habitat variation in fish assemblage  
218 structure over time was assessed using habitat type, time-period, and their interaction fitted as  
219 fixed effects, site as a random effect, and planned comparisons used post hoc to identify  
220 where changes occurred. Second, among-habitat variation in assemblage structure was  
221 assessed in each time-period using habitat (fixed effect), site (random effect), and Tukey  
222 multiple comparisons *post hoc* to identify where differences occurred (with *multcomp*). Count  
223 data models of fish species richness and total fish abundance were fit with Poisson, and

224 negative binomial distributions respectively, to accommodate non-stable variances and  
225 alternative exponential residual distributions (with glmer in *lme4*). Continuous measures of  
226 fish functional richness, functional dispersion, functional originality, and the CWM values for  
227 all four PCoA axes were normally distributed and were fitted with a Gaussian residual  
228 structure (with lme in *nlme*). Functional originality was fitted with a constant variance  
229 structure to allow for existing heterogeneity of variance. Inter-habitat variation in the  
230 taxonomic composition of fish assemblages was tested using two-way nested  
231 PERMANOVAs for each period (maximum permutations = 9999) on square root transformed  
232 data, with habitat (fixed factor) and site (random factor). Due to limited available  
233 permutations, Monte Carlo sampling was used to estimate *post hoc* pairwise comparisons  
234 (Anderson & Robinson, 2003).

235

236 The relative influence of coral loss on variation in fish species richness, total abundance,  
237 functional richness, functional dispersion and functional originality was estimated using  
238 mixed effects multiple linear regression (total coral cover %, time-period, habitat, and period-  
239 habitat interaction as fixed effects, and site as a random effect). Information-theoretic model  
240 selection based on AICc ranking ( $\Delta AICc$  values  $< 2$ ) was used to quantify model uncertainty  
241 and the relative importance of each explanatory variable (all predictors had Spearman's  
242 correlation coefficients  $< 0.4$  and variance inflation factors,  $G\sqrt{VIF^{(1/(2*df))}} < 3$ ) (Burnham &  
243 Anderson, 2002). Models within  $\Delta AICc < 2$  with the fewest degrees of freedom were  
244 considered the most parsimonious.

245

246 Exploratory graphical analyses of residuals were used to confirm that assumptions of  
247 homogeneity of variance, normality, and independence were met for all linear regression  
248 modelling, and generalised mixed effects models fit with Poisson or negative binomial

249 distributions were tested for overdispersion. Multivariate analyses of benthic composition  
250 were performed in Primer v6 with +PERMANOVA add on package (Clarke & Warwick,  
251 2001). All other analyses were performed in R (R Development Core Team 2016), and the  
252 packages *ape*, *cluster*, *geometry*, *polycor*, *rcdd*, *vegan*, and the function *FDchange* in *FD*.

253

## 254 **Results**

255

### 256 Benthic composition

257

258 Analyses of benthic composition across the 16 study sites revealed six major habitat  
259 groupings characterised by a disproportionate cover (25–54% of the total benthos) of (i)  
260 *Porites cylindrica* (hereafter branching *Porites*), (ii) massive *Porites* (mostly *Porites lutea*),  
261 (iii) *Pocillopora damicornis* (hereafter *Pocillopora*), (iv) soft coral, (v) mixed coral  
262 assemblages, and (vi) dead coral and macroalgae (<10% coral cover, hereafter degraded)  
263 (Fig. S1; PERMANOVA: Pseudo- $F = 8.09$ ,  $P = 0.0001$ , all pairwise comparisons  $P \leq 0.0002$ ;  
264 Table S2 and S3). These ‘degraded’ sites were subject to localised disturbance from recent  
265 cyclones and predation by crown-of-thorns starfish that caused acute coral loss on those reefs.  
266 Four of the six broad habitat groupings were represented by three sites, while massive *Porites*  
267 and *Pocillopora* habitats were represented by two and one site(s), respectively. Prior to the  
268 bleaching event, total coral cover (hard and soft coral) was similar among surveyed coral  
269 habitats (mean  $\pm$  SE:  $50.4 \pm 3.6$ ), which all had significantly greater cover of live coral than  
270 degraded habitats (Tukey, all  $P < 0.002$ ). Coral bleaching was widespread across surveyed  
271 sites in April 2016 affecting 51% of total coral cover (31 out of 36 coral taxa; Table S4), and  
272 did not vary significantly among habitats (percentage bleached of total coral cover; all  
273 pairwise comparisons,  $P > 0.05$ ) (Table 1). However, the mortality of corals following the

274 bleaching event varied among taxa (primarily soft coral, branching *Porites* and *Acropora*;  
 275 Fig. S2), causing significant declines in coral cover by October 2016 in just three habitats:  
 276 soft coral (31%), mixed coral (18%), and branching *Porites* (15%) habitats (Table 2; Fig. 1;  
 277 Table S5). Coral cover remained higher in all coral habitats relative to the degraded habitat in  
 278 October 2016 (Tukey, all  $P < 0.01$ ).

279

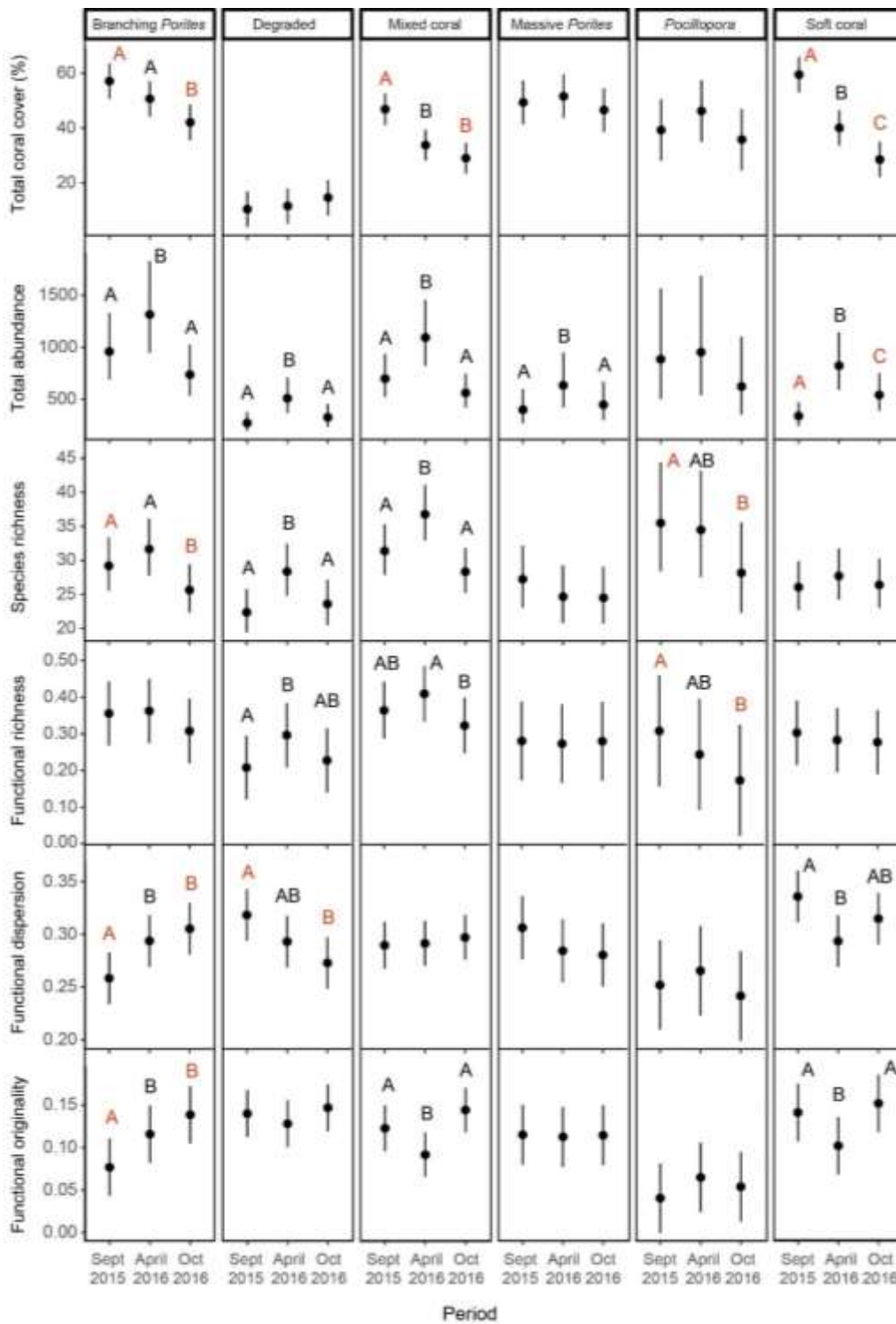
280 **Table 1** Mean  $\pm$  SE percent cover of benthic composition and coral bleaching extent within  
 281 habitats from September 2015 to October 2016.

Habitat	September 2015	April 2016			October 2016	Change in coral cover (Sept. 2015–Oct. 2016)
	Total coral cover	Total bleached coral cover	Percent bleached of total coral cover	Percent bleached of dominant coral taxa	Total coral cover	
Branching <i>Porites</i>	57.13 $\pm$ 2.79	21.81 $\pm$ 1.80	43.73 $\pm$ 2.92	45.86 $\pm$ 4.39	42.08 $\pm$ 2.30	-15.01 $\pm$ 3.39
Degraded	10.46 $\pm$ 1.80	7.04 $\pm$ 0.96	59.80 $\pm$ 4.74	-	14.63 $\pm$ 1.87	4.12 $\pm$ 3.39
Massive <i>Porites</i>	49.37 $\pm$ 2.81	23.82 $\pm$ 1.55	47.61 $\pm$ 4.06	46.96 $\pm$ 4.58	46.60 $\pm$ 3.19	-2.78 $\pm$ 4.15
Mixed coral	46.63 $\pm$ 3.07	15.03 $\pm$ 1.65	45.72 $\pm$ 4.18	-	29.03 $\pm$ 2.43	-17.88 $\pm$ 3.05
<i>Pocillopora</i>	39.31 $\pm$ 6.01	25.69 $\pm$ 1.73	56.68 $\pm$ 4.64	57.86 $\pm$ 4.58	35.83 $\pm$ 3.41	-3.47 $\pm$ 5.87
Soft coral	59.49 $\pm$ 1.93	27.27 $\pm$ 2.32	67.15 $\pm$ 3.23	67.95 $\pm$ 3.58	28.56 $\pm$ 2.65	-30.93 $\pm$ 3.39

282

283 Due to differential coral loss, coral cover varied among coral habitats following the  
 284 bleaching, with cover in mixed coral, and soft coral habitats significantly lower than  
 285 branching- and massive- *Porites* habitats, and *Pocillopora* habitat remaining intermediate  
 286 (Tukey, all  $P < 0.02$ ).

287



288

289 **Fig. 1** Within-habitat variation (fitted values  $\pm$  95% confidence intervals) in total coral cover  
 290 (%), fish species richness, total fish abundance, fish functional richness, functional dispersion,  
 291 and functional originality over survey periods. Contrasting letters indicate significant planned  
 292 comparisons within habitats over time (red letters highlight differences between September  
 293 2015 and October 2016).

294

295 Fish assemblage structure

296

297 The four PCoA axes cumulatively explained 56.8% of the projected inertia in the distribution

298 of fish species traits (Fig. 2 and 3). Generally, fish body-size and mobility increased along

299 PCoA 1 and 2, with diet groupings positioned along those gradients, ranging from small-

300 bodied site attached farming species through to larger, more mobile, piscivorous fish species

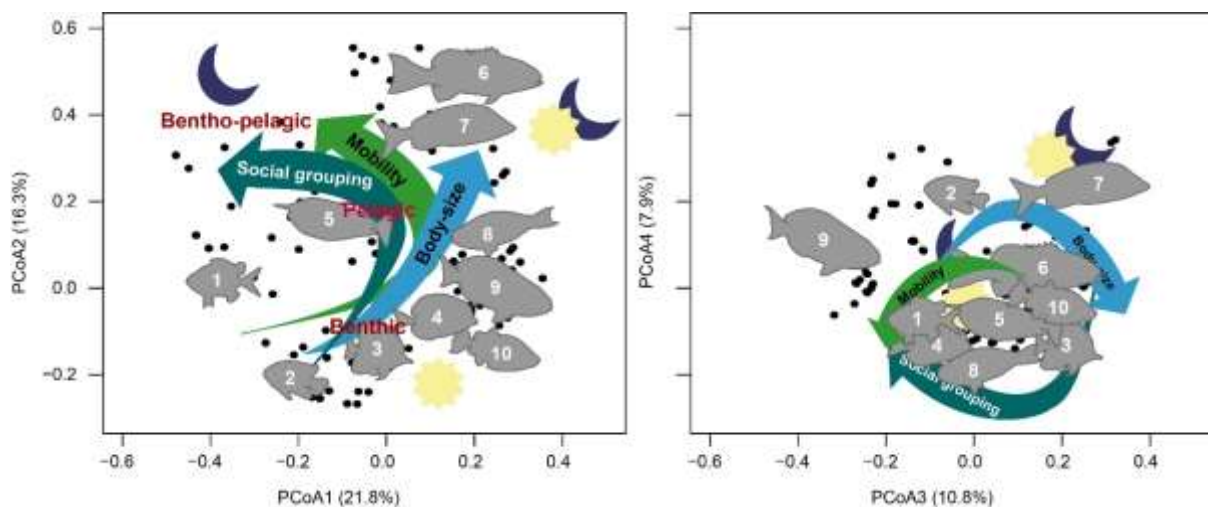
301 (Fig. 2). Nocturnally active, schooling planktivores occupied the left of PCoA1, and browsers

302 were positioned in the centre of PCoA 1 and 2. Generally, species clustered more closely on

303 PCoA 3 and 4. However, PCoA 4 separated excavating (PCoA3 left), farming (PCoA3

304 centre), and mixed diet feeding species (PCoA3 right), from all other species (Fig. 2).

305



306

307 **Fig. 2** Trait-based principal coordinates analysis of surveyed fish in functional space. Two

308 hundred and seventeen surveyed fish species (black dots) plotted four dimensions of functional

309 space defined by six traits: mean observed body-size (blue arrow indicating increasing size);

310 diet (fish symbols: 1, planktivore; 2, farmer; 3, corallivore; 4, grazer/detritivore; 5, browser; 6,

311 piscivore; 7, mixed diet; 8, scraper; 9, excavator; 10, invertivore); mobility (green arrow

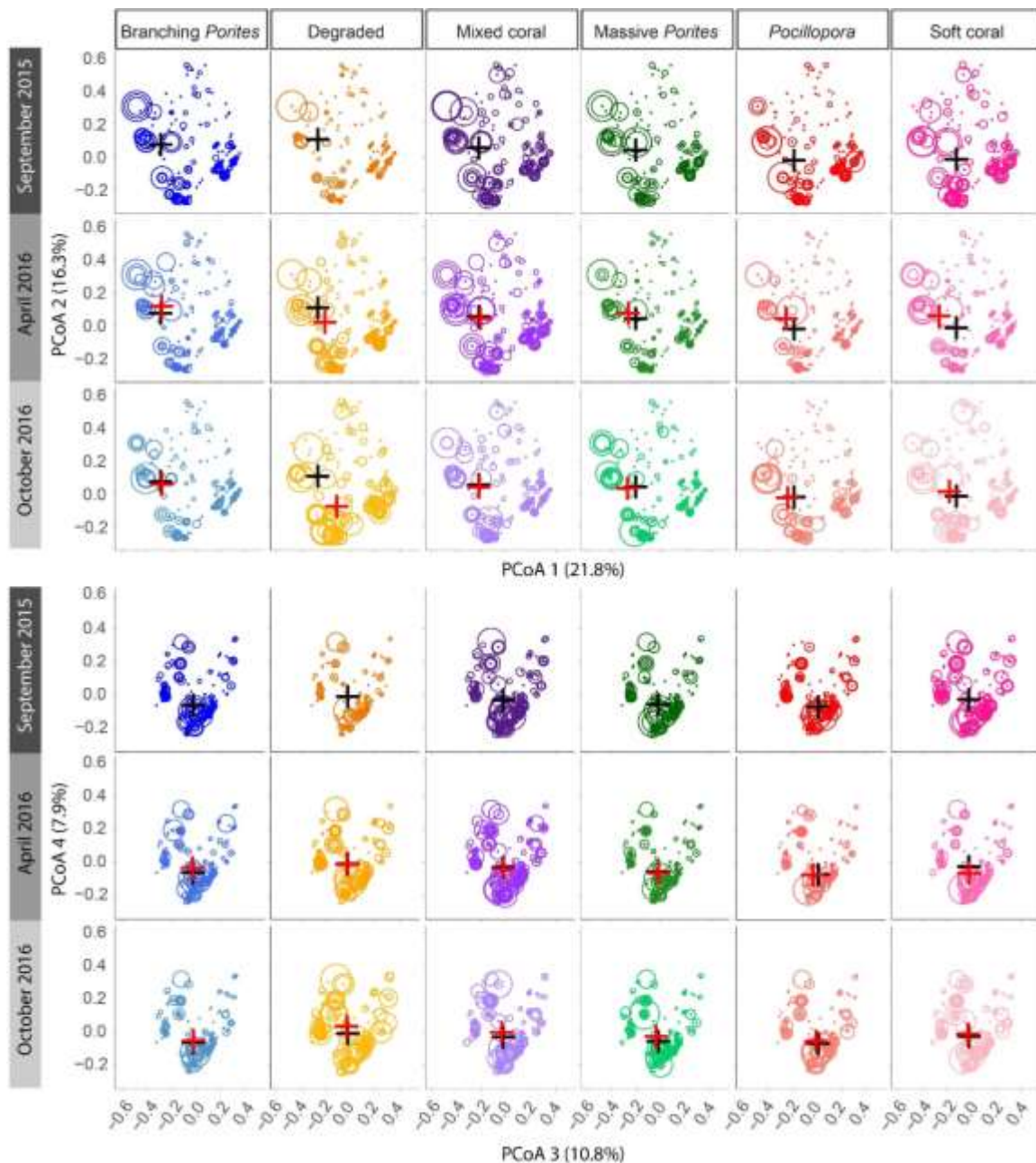
312 indicating increasing mobility); social grouping (dark blue arrow indicating gradient from

313 solitary fishes to large schools); time of activity (sun and/or moon); and position in the water  
314 column (red text). Illustrations show the average position of traits in the functional space.

315

316 Analysis of within-habitat variation in fish assemblage structure revealed changes in  
317 taxonomic (species richness and total abundance) and functional (functional richness,  
318 dispersion, and originality) characteristics in response to the bleaching event (Fig. 1 and 3).  
319 However, the nature of these changes varied among habitats (Table 2; Fig. 1; Table S5 and  
320 S6). Between September 2015 and October 2016, fish species richness declined in branching  
321 *Porites* and *Pocillopora* habitats; total fish abundance increased in soft coral habitats;  
322 functional richness declined in *Pocillopora* habitat; functional dispersion increased in  
323 branching *Porites* habitats, and decreased in degraded habitats; and functional originality  
324 increased in branching *Porites* habitats (Fig. 1 and 3).

325



326

327 **Fig. 3** Principal coordinates analyses (PCoA) of fish assemblage functional structure in habitats  
 328 in each period, showing the first two (top: PCoA 1 and 2), and second two (bottom: PCoA 3  
 329 and 4), dimensions of functional trait space. Circle sizes are proportional to species mean  
 330 relative abundances. Crosses mark community weighted mean centroids in each period (black:  
 331 September 2015; red: April 2016, or October 2016).

332



333 **Table 2** Significant planned comparisons (contrast effect size and 95% confidence intervals;  
 334 CI) of within-habitat variation in total coral cover and fish assemblage responses between  
 335 September 2015 and October 2016 among habitats.

336

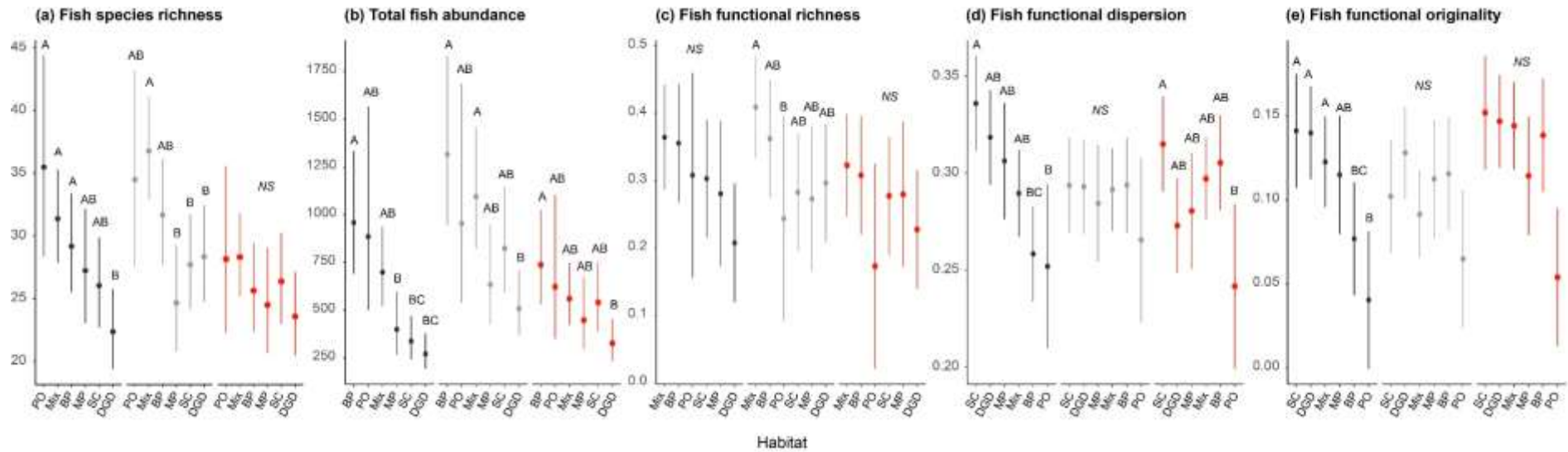
Response	Habitat	Period contrast	Lower CI	Upper CI	Test stat	df	P
Total coral cover	Branching <i>Porites</i>	-15.049	-21.725	-8.373	-4.44	265	<0.001
	Mixed coral	-17.875	-23.882	-11.867	-5.86	265	<0.001
	Soft coral	-30.927	-37.603	-24.251	-9.12	265	<0.001
Fish species richness	Branching <i>Porites</i>	0.879	0.775	0.996	-2.03	266	0.04
	<i>Pocillopora</i>	0.793	0.648	0.972	-2.25	266	0.02
Total fish abundance	Soft coral	1.597	1.210	2.108	3.32	265	<0.001
Functional richness	<i>Pocillopora</i>	-0.135	-0.255	-0.015	-2.22	265	0.03
Functional dispersion	Branching <i>Porites</i>	0.047	0.018	0.077	3.14	265	0.002
	Degraded	-0.046	-0.075	-0.016	-3.04	265	0.003
Functional originality	Branching <i>Porites</i>	0.062	0.024	0.099	3.24	260	0.001

337

338

339 These changes caused an increased overall similarity in the taxonomic and functional  
 340 diversity of fish assemblages across distinct habitat types. Analysis of among habitat  
 341 variation in fish assemblage structure over time revealed that prior to the bleaching event  
 342 there was significant variation in species richness, total fish abundance, fish functional  
 343 dispersion, and functional originality, with habitat type accounting for 14–39% of the  
 344 variation in these indices (not including fish abundance). Eight months after the bleaching  
 345 event, however, this variation was almost entirely lost, with habitat type accounting for just  
 346 8–17%) of the variation in those measures (Table 3; Fig. 4). There was also an increased  
 347 similarity in the taxonomic and functional composition of fish assemblages after the  
 348 bleaching event. The number of pairwise differences in taxonomic composition among  
 349 habitats declined from 7 in September 2015, to 5 in April 2016, and just 4 by October 2016  
 350 (PERMANOVAs: September 2015: Pseudo- $F = 2.42$ ,  $df = 5$ ,  $P = 0.0001$ , all pairwise  
 351 comparisons  $P < 0.05$ ; April 2016: Pseudo- $F = 2.02$ ,  $df = 5$ ,  $P = 0.001$ , all pairwise  
 352 comparisons  $P < 0.05$ ; October 2016: Pseudo- $F = 1.94$ ,  $df = 5$ ,  $P = 0.001$ , all pairwise  
 353 comparisons  $P < 0.03$ ). Similarly, the number of pairwise differences in functional

354 composition indicated by CWMs across the four PCoA axes declined from 11 in September  
355 2015, to two in April 2016, and 4 by October 2016 (Table 4). Changes in CWM for PCoA  
356 axes 1 and 2 indicated no changes in dominant traits across habitats though time. However,  
357 CWM values on PCoA 3 and 4 exhibited significant directional shifts overall between  
358 September 2015 and October 2016, indicated by back (September 2015) and red (October  
359 2016) centroid cross markers, towards small-bodied habitat generalist algae-farming species  
360 (Table 4; Fig. 2 and 3).



361

362 **Fig. 4 Among-habitat variation in** fish assemblage structure within each time-period (fitted values  $\pm$  95% confidence intervals): September 2015  
 363 (black); April 2016 (grey), and; October 2016 (red). Different letters illustrate significant differences among habitats within each period revealed  
 364 by linear mixed effects models (habitat as fixed effect) and Tukey pair-wise comparisons *post hoc* ( $P < 0.05$ ). Habitats: BP, branching *Porites*;  
 365 DGD, degraded reef; Mix, mixed coral; MP, massive *Porites*; PO, *Pocillopora*; and SC, soft coral.

366 **Table 3** Components of variation for mixed effects models of fish assemblage structure among  
 367 habitats in each survey period and Tukey multiple comparisons (number of among-habitat  
 368 differences in brackets shown in bold). Habitats: BP, branching *Porites*; DGD, degraded; Mix,  
 369 mixed coral; MP, massive *Porites*; PO, *Pocillopora*; SC, soft coral.  
 370

Fish response metric	Period	$R^2$ fixed (habitat)	$R^2$ random (site)	$R^2$ residuals (transects)	Habitat differences ( $P < 0.05$ )
Species richness	Sept-15	0.28	0.11	0.61	<b>(3)</b> DGD $\neq$ BP, Mix, PO
	April-16	0.32	0.13	0.55	<b>(3)</b> Mix $\neq$ DGD, MP, SC
	Oct-16	0.08	0.22	0.69	<b>(0)</b>
Total fish abundance †	Sept-15				<b>(7)</b> BP $\neq$ DGD, MP, SC; Mix, PO $\neq$ DGD, SC
	April-16				<b>(2)</b> BP, Mix $\neq$ DGD
	Oct-16				<b>(1)</b> BP $\neq$ DGD
Functional richness	Sept-15	0.14	0.31	0.55	<b>(0)</b>
	April-16	0.23	0.13	0.64	<b>(1)</b> SC $\neq$ Mix
	Oct-16	0.10	0.26	0.64	<b>(0)</b>
Functional dispersion	Sept-15	0.33	0.10	0.57	<b>(3)</b> BP $\neq$ DGD, SC; SC $\neq$ PO
	April-16	0.02	0.09	0.89	<b>(0)</b>
	Oct-16	0.14	0.02	0.84	<b>(1)</b> SC $\neq$ PO
Functional originality	Sept-15	0.39	0.03	0.59	<b>(7)</b> BP, PO $\neq$ DGD, Mix, SC; PO $\neq$ MP
	April-16	0.04	0.05	0.91	<b>(0)</b>
	Oct-16	0.15	0.24	0.60	<b>(0)</b>

371 †  $R^2$  estimations not available for generalized linear mixed models with negative binomial distribution.

372

373

374

375 **Table 4** Variation in community abundance-weighted means on each PCoA axes indicating  
 376 change in the functional structure of fish assemblages through time, and differences among  
 377 habitats (Tukey,  $P < 0.05$ ). The number of inter-habitat differences within each period are  
 378 shown for each axis.

PCoA axis	PCoA axis inertia explained (%)	Change (Sept 2015 – Oct 2016)	Overall habitat differences	Habitat differences ( $P < 0.05$ )		
				Sept 2015	April 2016	Oct 2016
PCoA 1	21.8	No	BP $\neq$ DGD, Mix, SC	2	0	2
PCoA 2	16.3	No	BP $\neq$ DGD, Mix	0	2	1
PCoA 3	10.8	Yes	PO $\neq$ DGD	5	0	1
PCoA 4	7.9	Yes	n/a	4	0	0

379  
 380 Changes in fish assemblage structure across habitats between September 2015 and October  
 381 2016 were largely unrelated to levels of coral loss (Table 2; Fig. 1; Table S5). For example,  
 382 in the three habitats that experienced significant coral loss, there were either no detected  
 383 changes in fish assemblages (mixed coral); total abundance increased (soft coral); or species  
 384 richness declined, and functional dispersion and functional originality increased (branching  
 385 *Porites*) (Fig. 1; Table S6). Additionally, species- and functional richness declined in  
 386 *Pocillopora* habitat, and fish functional dispersion increased in degraded habitats despite no  
 387 change in coral cover (Table 2; Fig. 1). Analyses of the relative influence of coral loss on  
 388 variation in all five measures of fish assemblage structure revealed total coral cover in all top  
 389 candidate multiple linear regression models. However it was not the best predictor of  
 390 variation in any metric (species richness, total abundance, functional richness, functional  
 391 dispersion and functional originality) relative to the influence of time-period or habitat type  
 392 (or their interaction; Table S7).

393

## 394 Discussion

395

396 Our study of the impacts of mass coral bleaching on coral reef fish assemblage structure  
397 across six taxonomically distinct reef habitats revealed three key findings. First, we show that  
398 bleaching disturbance increased the taxonomic and functional similarity of fish assemblage  
399 structure due to species turnover among previously differentiated fish assemblages (i.e.,  
400 biotic homogenization). Second, we found that despite limited change in taxonomic or  
401 functional richness across all habitats, the functional originality of fish assemblages  
402 increased. This is potentially important because it suggests that there was a net loss of  
403 functional redundancy at the system level. Third, we found that pre-bleaching coral species  
404 composition was a better predictor of changes in fish assemblage structure, than absolute  
405 coral loss. These results emphasize how measures of alpha diversity (taxonomic or functional  
406 richness) can mask important changes in ecosystems as assemblages reorganize with  
407 implications for ecosystem function. Our findings also highlight the important but poorly  
408 understood role of coral species composition in structuring communities and influencing  
409 cross-scale resilience to system-wide disturbance due to the response diversity of corals and  
410 reef fishes.

411

412 The system-wide biotic homogenization among previously differentiated reef fish  
413 assemblages was evident by the increased similarity in the taxonomic and functional  
414 composition of fish assemblages following the bleaching event. Prior habitat-related  
415 differences in fish species richness, total abundance, functional dispersion, functional  
416 originality, and the taxonomic and functional composition of fish assemblages was largely  
417 lost after the bleaching event, despite no change in the taxonomic or functional richness  
418 across all surveyed reefs (local alpha diversity). Biotic homogenization is increasingly  
419 considered a central component of the broader biodiversity crisis (Olden et al., 2004;  
420 McKinney & Lockwood, 1999, McGill et al., 2015). Whilst disturbance-mediated loss of

421 species has occurred across ecosystems (Olden et al., 2004; Butchart et al., 2010), meta-  
422 analyses show no systematic loss of species richness, rather a loss of beta diversity,  
423 suggestive of increasing biotic homogenization at a global scale (Dornelas et al., 2014).  
424 Biotic homogenization has been documented across several groups including temperate  
425 marine fishes (Magurran, Dornelas, Moyes, Gotelli, & McGill, 2015), freshwater fishes  
426 (Villéger, Grenouillet, & Brosse, 2014), birds (Cassey, Lockwood, Blackburn, & Olden,  
427 2007), and plants (Schwartz, Thorne, & Viers, 2006; Rooney, 2009). We add to these studies  
428 with evidence of a high diversity coral reef ecosystem response to climatic disturbance,  
429 showing that isolated measures of  $\alpha$  diversity (i.e. richness) can conceal important spatial beta  
430 diversity trends, such that the presence of species does not capture the rapid reorganisation of  
431 assemblages (Magurran et al., 2015; McGill et al., 2015).

432

433 The consequences of biotic homogenization are not yet well understood (Olden et al., 2004).  
434 However, trait-based functional approaches are increasingly used to provide insight into  
435 ecological changes that may influence the functional stability of ecosystems (Mouillot et al.,  
436 2013; Graham, Jennings, MacNeil, Mouillot, & Wilson, 2015). We observed an increased  
437 similarity of functional dispersion (trait variability), functional originality (trait uniqueness),  
438 and functional composition of reef fish assemblages following the bleaching event.  
439 Furthermore, we found subtle but significant directional shifts in dominant fish traits across  
440 habitats towards algae-farming, small-bodied habitat generalists, causing an increased  
441 homogenization of trait space. These findings are consistent with reports of functional  
442 homogenization occurring across scales and ecosystems (Devictor, Julliard, Couvet, Lee, &  
443 Jiguet, 2007; Abadie, Machon, Muratet, & Porcher, 2011; Pool & Olden, 2012; Villéger et  
444 al., 2014). The susceptibility of a species to extirpation is typically non-random, and related  
445 to organism traits such as body-size, diet, fecundity, growth-rate, habitat specialisation, social



446 grouping and abundance (McKinney, 1997; Graham et al., 2011). As predicted by niche  
447 theory (Hutchinson, 1957), building evidence suggests that global changes are causing  
448 worldwide declines of specialist species which have effectively been replaced by generalists  
449 thereby increasing functional similarity at the community level (Clavel et al., 2011). Indeed,  
450 severe coral bleaching events tend to favour larger-bodied, habitat generalists at the expense  
451 of obligate coral feeders and small-bodied fishes reliant on live coral habitat structure  
452 (Wilson et al., 2006; Pratchett et al., 2011). However, trait vulnerability to bleaching events  
453 can vary with the rate and extent that coral tissue mortality also leads to the loss of the  
454 physical coral structures (Wilson et al., 2006). Within the current study, coral tissue mortality  
455 caused a loss of soft coral colony structures (Ferrari 2017), but not of hard corals whose  
456 carbonate colony structures remained largely intact by October 2016. Experimental evidence  
457 suggests that soft coral exerts little direct influence on reef fish assemblages (Lewis 1998).  
458 Therefore, shifts in dominant traits from habitat specialists to generalists appear likely to  
459 result from the loss of coral tissue rather than habitat structural complexity.

460

461 Together with taxonomic and functional homogenization, we detected an increase in  
462 functional originality at the system level, inferring overall loss of functional redundancy of  
463 fish assemblages following coral bleaching, and providing further evidence of non-random  
464 trait filtering. The increase in functional originality was largely driven by species shifts in  
465 branching *Porites* habitats, where live coral tissue was lost though colony structures remained  
466 intact, and fishes with a strong preference for live branching coral (e.g. small-bodied, site-  
467 attached, schooling planktivores), were replaced by a range of other functional entities  
468 including algae-farming, habitat generalists. As habitat disturbance reduces the abundance of  
469 functionally similar species, the number of species contributing to any one function (i.e.,  
470 functional redundancy) may decline as remaining species become relatively more

471 functionally unique (Walker, 1992; Reich et al., 2012). We found that trait-filtering of habitat  
472 specialists likely reduced the competitive exclusion of other functional entities, facilitating  
473 habitat use by a range of other groups and reducing functional redundancy at the community  
474 level (MacArthur & Levins, 1967). We provide clear data showing that whilst functional  
475 richness remained consistent across the bleaching event, the turnover of fish species altered  
476 the distribution of traits in specific habitats. However, changes were poorly predicted by the  
477 loss of coral at the system level.

478

479 Interestingly, pre-bleaching coral species composition was a better predictor of changes in  
480 fish assemblage structure following the bleaching event than the specific level of coral loss  
481 within each habitat type. Whilst substantial coral loss typically incurs concurrent declines in  
482 the species richness and total abundance of reef fishes after bleaching (>20% total cover and  
483 reduction in absolute cover below 10%; Wilson et al., 2006), we show that observed changes  
484 in fish assemblage structure (including species richness and total abundance) were variable  
485 among habitats and inconsistent with changes in coral cover. For example, the greatest  
486 changes in fish assemblage structure occurred in branching *Porites* habitats that lost 15%  
487 coral cover. However, in mixed and soft coral habitats that experienced greater coral loss  
488 (18% and 31%, respectively), fish assemblage structure only changed in soft coral habitats  
489 (the only habitat where the physical coral structures were also lost) where total fish  
490 abundance increased. Further, changes in fish assemblage structure were evident in habitats  
491 where significant coral loss was not detected (e.g. reduced taxonomic and functional richness  
492 in *Pocillopora* habitat). With widespread coral bleaching across habitats, our results illustrate  
493 the differential susceptibility of distinct reef fish assemblages, governed primarily by the  
494 vulnerability of particular fish traits associated with specific coral configurations. Coral  
495 habitats provide a range of niche spaces for the coexistence of diverse but functionally

496 unbalanced fish assemblages (Mouillot et al., 2014; Brandl, Emslie, Ceccarelli, & Richards,  
497 2016; Richardson, Graham, & Hoey, 2017). Some provide particular niche habitat to  
498 specialist guilds of fishes (Richardson, Graham, Pratchett, et al., 2017), such that relatively  
499 small habitat disturbance may incur disproportionate change in fish assemblage structure.  
500 Indeed, we found that fish assemblages in branching coral habitats (branching *Porites* and  
501 *Pocillopora*) were most negatively impacted by the bleaching event, despite differential coral  
502 loss. Conversely, coral reefs with high or diverse initial coral cover that incur coral loss may  
503 have limited effects on fish populations if the availability of required habitat remains  
504 sufficient (e.g. corallivores, Pratchett, Wilson, & Baird, 2006), such as in mixed coral habitats  
505 in our study.

506

507 These results provide some insight into the underlying dynamics of coral bleaching on reef  
508 fish assemblages, and highlight the role of coral species composition in determining the  
509 susceptibility of assemblage structure to disturbance. However, further examination of coral  
510 habitats across locations, particularly reefs characterised by dominant though thermally  
511 sensitive *Acropora*, would enable better understanding of how shifts in coral species  
512 composition may affect changes in fish assemblage structure at a broader scale. Furthermore,  
513 our results only provide a snapshot of a highly complex, dynamic system, and caution is  
514 recommended for applying these findings to longer-term predictions. It is unknown if the  
515 observed functional trait shifts constitute a temporary phase in the reassembly of coral reefs  
516 directly following acute bleaching disturbance, or if shifts will be sustained (Bellwood et al.,  
517 2012). Similarly, lag-responses to bleaching events have been observed in corals (e.g.  
518 disease; Bruno et al., 2007; Miller et al., 2009), and populations of coral reef fishes where the  
519 delayed erosion of dead coral structures has caused substantial declines in smaller size-  
520 classes of reef fish (Wilson et al., 2006; Graham et al., 2007). There is much uncertainty

521 about how distinct and emerging configurations of species will respond to ongoing  
522 disturbances and how important ecosystem processes will be maintained (Graham et al.,  
523 2014). A great deal more research is needed in this area to understand reef dynamics in the  
524 Anthropocene, so to guide viable and pragmatic management approaches (Norström et al.,  
525 2016). Finally, whilst trait-based estimations of ecological diversity provide a useful tool for  
526 assessing the functional trajectory of disturbed ecosystems, we acknowledge that they remain  
527 coarse approximations based on our current understanding of species' roles that do not likely  
528 capture finer-scale, nuanced niche partitioning (McGill, Enquist, Weiher, & Westoby, 2006).  
529  
530 Climate induced mass coral bleaching often devastates coral reef ecosystems, threatening the  
531 livelihoods of people that rely upon the ecosystem services that reefs provide (Moberg &  
532 Folke, 1999; Norström et al., 2016). As global temperatures rise, new governance challenges  
533 are faced by managers as non-random species shifts transform reef assemblages into novel  
534 configurations of species (Graham et al., 2014), with unknown implications for their capacity  
535 to maintain key ecosystem functions (Norström et al., 2016; Hughes, Barnes, et al., 2017).  
536 Comparing the impacts of bleaching on reef fish assemblage structure across distinct reef  
537 habitats provided some insight into the variable responses of both corals and fishes to thermal  
538 stress, and highlighted the potential homogenization of fish assemblages across scales. Biotic  
539 homogenization is increasingly considered one of the most pressing, but largely  
540 unrecognized, biodiversity crises faced globally (Dornelas et al., 2014; Magurran et al., 2015;  
541 McGill et al., 2015). Our results underscore the importance of coral species composition in  
542 determining ecological organisation and the susceptibility of reef ecosystems to disturbance.  
543 As species shifts persist, the suitability of coral reef management and the value of different  
544 coral dominated states will likely hinge upon the capacity of those ecosystems to maintain  
545 key ecological processes with ongoing disturbance.

546

547 **Acknowledgements**

548

549 We thank Lizard Island Research Station staff for field support, Valeriano Parravicini for  
550 providing trait information for some species, Murray Logan and Rhondda Jones for statistical  
551 advice, and Michael McWilliam and two anonymous reviewers for useful comments. This  
552 study was funded by the Australian Research Council (ARC) to ASH (DE130100688) and  
553 NAJG (DE130101705), and the ARC Centre of Excellence for Coral Reef Studies to LER  
554 (Higher Degree Research Enhancement Scheme).

555

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